

CRF Errors Corrected by the STIC Systems Branch

CRF Processing Date: 12/4/02
 Edited by: DC (STIC staff)
 Verified by: W/H

Serial Number: 091171671

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: **REMOVED**
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error _____)
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error _____)

Examiner: _____ The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form



PCT09

RAW SEQUENCE LISTING

PATENT APPLICATION N: US/09/171,671

DATE: 11-14-01

TIME: 11:11:14

fk

Input File : A:\PTO.DC.txt

Output File : N:\CRF4\12032002\I171671.raw

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RAW SEQUENCE LISTING

PATENT ABSTRACT N: US/09/171,671

DATE: 11-04-01

TIME: 14:14

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Output File: N:\CRF4\12032002\I171671.raw

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W--> 75 Val Ala Xaa Ser Tyr Asp

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W--> 97 Ala Xaa Ser Tyr Asp

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RAW SEQUENCE LISTING

PATENT ABSTRACT N: US/09/171,671

DATE: 11-04-2009

TIME: 11:14:14

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Output File: N:\CRF4\12032002\I171671.raw

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W--> 207 Val Ala Xaa Ser Tyr

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RAW SEQUENCE LISTING

PATENT ABSTRACT N: US/09/171,671

DATE: 11-04-2009

TIME: 14:14:14

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File Name: N:\CRF4\12032002\I171671.raw

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[illegible]

Figure 1. Schematic representation of the experimental design. The subjects were divided into two groups: the control group and the experimental group. The control group was divided into two subgroups: the control group and the control group. The experimental group was divided into two subgroups: the experimental group and the experimental group.

File Name : A:\PTO.DC.txt

File Name: N:\CRF4\12032002\I171671.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

[illegible]

RAW SEQUENCE LISTING ERROR SUMMARY

US/09/171,671

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1. $\frac{1}{2} \times \frac{1}{2} = \frac{1}{4}$

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[illegible]

VERIFICATION SUMMARY

US/09/171,671

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Figure 1. The effect of the concentration of the *Agrobacterium* suspension on the transformation efficiency of *Agrobacterium* strains. The number of transformed cells was determined by the number of colonies obtained on the selective medium. The results are the mean of three independent experiments. Error bars represent the standard deviation.

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[illegible]

VERIFICATION SUMMARY

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DATE: 11/14/02

TIME: 11:14:14

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Output File: N:\CRF4\12032002\I171671.raw

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